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## **EUROPEAN PATENT APPLICATION**

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#### Remarks:

The applicant has subsequently filed a sequence listing and declared, that it includes no new matter.

(54) Survival motor neuron (SMN) gene: a gene for spinal muscular atrophy

(57) The present invention relates to the discovery of a survival motor-neuron gene or SMN gene which is a chromosome 5-SMA (Spinal Muscular Atrophy) determining gene. The present invention further relates to the nucleotide sequence encoding the SMN gene and corresponding amino acid sequence, a vector containing the gene encoding the SMN protein or a DNA sequence corresponding to the gene and transformant strains containing the SMN gene or a DNA sequence corresponding to the gene.

The present invention also relates to means and methods for detecting motor neuron diseases having symptoms of muscular weakness with or without sensory changes such as amytrophic lateral sclerosis (ALS), spinal muscular atrophy (SMA), primary lateral sclerosis (PLS) and the like.

MAMSSGGSGGVPEQEDSVLFRRGTGQSDDSDIWDDTALIKAYDKAVAS

FKHALKNGDICETSGKPKTTPKRKPAKKNKSQKKNTAASLQQWKVGDKCSAIWSEDGCIY

PATIASIDFKRETCVVVYTGYGNREEQNLSDLLSPICEVANNIEQNAQENENESQVSTDE

SENSRSPGNKSDNIKPKSAPWNSFLPPPPPPMPGPRLGPGKPGLKFNGPPPPPPPPPHLL

SCWLPPFPSGPPIIPPPPPICPDSLDDADALGSMLISWYMSGYHTGYYMGFRQNQKEGRC

SHSLN

CGGGGCCCACGCTGCGCACCCGCGGTTTGCTATGGCGATGAGCAGCGGCGGCAGTGGT GATTCTGACATTTGGGATGATACAGCACTGATAAAAGCATATGATAAAGCTGTGGCTTCA TTTAAGCATGCTCTAAAGAATGGTGACATTTGTGAAACCTCGGGTAAACCAAAAACCACA CCTAAAAGAAACCTGCTAAGAAGAATAAAAGCCAAAAGAAGAATACTGCAGCTTCCTTA CAACAGTGGAAAGTTGGGGACAAATGTTCTGCCATTTGGTCAGAAGACGGTTGCATTTAC CCAGCTACCATTGCTTCAATTGATTTTAAGAGAGAAACCTGTGTTGTGGTTTACACTGGA **TATGGAAATAGAGAGGAGCAAAATCTGTCCGATCTACTTTCCCCAATCTGTGAAGTAGCT** AATAATATAGAACAGAATGCTCAAGAGAATGAAAATGAAAGCCAAGTTTCAACAGATGAA <u>AGTGAGAACTCCAGGTCTCCTGGAAATAAATCAGATAACATCAAGCCCAAATCTGCTCCA</u> TGGAACCCCTTTCTCCCTCCACCACCCCCATGCCAGGGCCAAGACTGGGACCAGGAAAG <u>TCATGCTGGCTGCCTCCATTTCCTTCTGGACCACCATAATTCCCCCCACCACCTCCCATA</u> **TGTCCAGATTCTCTTGATGATGCTGATGCTTTGGGAAGTATGTTAATTTCATGGTACATG** TCACATTCCTTAAATTAAGGAGAAATGCTGGCATAGAGCAGCACTAAATGACACCACTAA AGAAACGATCAGACAGATCTGGAATGTGAAGCGTTATAGAAGATAACTGGCCTCATTTCT TCAAAATATCAAGTGTTGGGAAAGAAAAAGGAAGTGGAATGGGTAACTCTTCTTGATTA **AAAGTTATGTAATAACCAAATGCAATGTGAAATATTTTACTGGACTCTTTTGAAAAACCA** TCTGTAAAAGACTGAGGTGGGGGGGGGGGGCAGCACGGTGGTGAGGCAGTTGAGAAAAT **AGAAGGGTGTTGTAGTTTATAAAAGACTGTCTTAATTTGCATACTTAAGCATTTAGGAAT** GTGGCAAAATGTTACAGAATCTAACTGGTGGACATGGCTGTTCATTGTACTGTTTTTTTC **KAAAAAAAAAAAAAA** 

FIGURE 2A

AATTTTTAAATTTTTTGTAGAGACAGGGTCTCATTATGTTGCCCAGGGTGGTGAAGCTCCA GGTCTCAAGTGATCCCCCTACCTCCGCCTCCCAAAGTTGTGGGATTGTAGGCATGAGCCACTG CAAGAAAACCTTAACTGCAGCCTAATAATTGTTTTCTTTGGGATAACTTTTAAAGTACATTAA AAGACTATCAACTTAATTTCTGATCATATTTTGTTGAATAAAATAAGTAAAAATGTCTTGTGAA TGCTCACATTCCTTAAATTAAGGA\*GTAAGTCTGCCAGCATTATGAAAGTGAATCJ"TACTTTT GTAAAACTTTATGGTTTGTGGAAAAACAATGTTTTTGAACAGTTAAAAGTTCAGATGTTA*G*A AAGTTGAAAGGTTAATGTAAAACAATCAATATTAÁAGAATTTTGATGCCAAAACTATTAGATA ATACTTTCACAATAAAGAGCTTTAGGATATGATGCCATTTTATATCACTAGTAGGCAGACCAG GAAGTGCTCTACTCAAGTTTAACTGGTGTCCACAGAGGGACATGGTTTAACTGGAATTCGTCAA GCCTCTGGTTCTAATTTCTCATTTGCAG\*GAAATGCTGGCATAGAGCAGCACTAAATGACACC ACTAAAGAAACGATCAGACAGATCTGCAATGTGAAGCGTTATAGAAGATAACTGGCCTCATTT <u>CTTCAAAATATCAAG</u>TGTTGGGAAAGAAAAAAGGAAGTGGAATGGGTAACTCTTCTTGATTA AAAGTTATGTAATAACCAAATGCAATGTGAAATATTTTACTGGACTCTTTTGAAAAAC CATCTGTAAAAGACTGGGGTGGGGGTGGGGGGGCAGCACGGTGGTGAGGCAGTTGAGAAAA **GAGAAGGGTGTTGTAGTTTATAAAAGACTGTCTTAATTTGCATACTTAAGCATTTAGG** ATGTGGCAAAATGTTACAGAATCTAACTGGTGGACATGGCTGTTCATTGTACTGTTTTTT 

FIGURE 2B

CGGGGCCCACGCTGCGCATCCGCGGGTTTGCTATGGCGATGAGCAGCGGCGGCAGTGGT <u>GGCGGCGTCCC</u>GGAGCAGGAGGATTCCGTGCTGTTCCGGCGCGCGCACAGGCCAG\*AGCGAT <u>GATTCTGACATTTGGGATGATACAGCACTGATAAAAGCATATGATAAAGCTGTGGCTTCA</u> TTTAAGCATGCTCTAAAGAATGGTGACATTTGTGAAACTTCGGGTAAACCAAAAACCACA CCTANANGANAACCTGCTAAGAAGAATAAAAGCCAAAAGAAGAATACTGCAGCTTCCTTA CAACAG\*TGGAAAGTTGGGGACAAATGTTCTGCCATTTGGTCAGAAGACGGTTGCATTTAC <u>CCAGCTACCATTGCTTCAATTGATTTTAAGAGAGAAACCTGTGTTGTGGTTTACACTGGA</u> <u>TATGGAAATAGAGAGGAGCAAAATCTGTCCGATCTACTTTCCCCAATCTGTGAAGTAGCT</u> AATAATATAGAACAGAATGCTCAAGAG\*AATGAAAATGAAAGCCAAGTTTCAACAGATGAA <u>AGTGAGA</u>ACTCCAGGTCTCCTGGAAATAAATCAGATAACATCAAGCCCAAATCTGCTCCA TGGAACTCTTTTCTCCCTCACCACCCCCATGCCAGGGCCAAGACTGGGACCAGGAAAG \*CCAGGTCTAAAATTCAATGGCCCACCACCACCGCCACCACCACCACCACCACTTACTA TCATGCTGGCTGCCTCCATTTCCTTCTGGACCACCA\*ATAATTCCCCCACCACCACCATA TGTCCAGATTCTCTTGATGATGCTGATGCTTTGGGAAGTATGTTAATTTCATGGTACATG AGTGGCTATCATACTGGCTATTATATG\*GGTTTCAGACAAAAATCAAAAAGAAGGAAGGTGC TCACATTCCTTAAATTAAGGA\*GAAATGCTGGCATAGAGCAGCACTAAATGACACCACTAA AGAAACGATCAGACAGATCTGGAATGTGAAGCGTTATAGAAGATAACTGGCCTCATTTCT TCAAAATATCAAGTGTTGGGAAAGAAAAAAGGAAGTGGAATGGGTAACTCTTCTTGATTA AAAGTTATGTAATAACCAAATGCAATGTGAAATATTTTACTGGACTCTTTTGAAAAAC CATCTGTAAAAGACTGGGGTGGGGGTGGGACGCCAGCACGGTGGTGAGGCAGTTGAGAAAA **GAGAAGGGTGTTGTAGTTTATAAAAGACTGTCTTAATTTGCATACTTAAGCATTTAGG ATGTGGCAAAATGTTACAGAATCTAACTGGTGGACATGGCTGTTCATTGTACTGTTTTTT Τ**CΓΑΓCΤΤCΤΑΓΑΤGTTTAAAAGTATAATAAAAATA ΤΑΤΤΑΑΤΥΤΤΤΤΤΤΑΑΑΑΑΑΑΑ 

FIGURE 3A

AATTTTTAAATTTTTTGTAGAGACAGGGTCTCATTATGTTGCCCAGGGTGGTGTCAAGCTCCA GGTCTCAAGTGATCCCCCTACCTCCGCCTCCCAAAGTTGTGGGATTGTAGGCATGAGCCACTG CAAGAAAACCTTAACTGCAGCCTAATAATTGTTTTCTTTGGGATAACTTTTAAAGTACATTAA **AAGACTATCAACTTAATTTCTGATCATATTTTTGTTGAATAAAATAAGTAAAATGTCTTGTGAA** TTTTTTTAACTTCCTTTATTTTCCTTACAG\*GGTTTCAGACAAAATCAAAAGAAGGAAGG TGCTCACATTCCTTAAATTAAGGA\*GTAAGTCTGCCAGCATTATGAAAGTGAATCTTACTTTT GTAAAACTTTATGGTTTGTGGAAAACAAATGTTTTTGAACAGTTAAAAAGTTCAGATGTTAAA AAGTTGAAAGGTTAATGTAAAACAATCAATATTAAAGAATTTTGATGCCAAAACTATTAGATA ATACTTTCACAATAAAGAGCTTTAGGATATGATGCCATTTTATATCACTAGTAGGCAGACCAG CAGACTTTTTTTTTTTTGTGATATGGGATAACCTAGGCATACTGCACTGTACACTCTGACATAT GAAGTGCTCTAGTCAAGTTTAACTGGTGTCCACAGAGGACATGGTTTAACTGGAATTCGTCAA GCCTCTGGTTCTAATTTCTCATTTGCAG\*GAAATGCTGGCATAGAGCAGCACTAAATGACACC ACTAAAGAAACGATCAGACAGATCTGGAATGTGAAGCGTTATAGAAGATAACTGGCCTCATTT CTTCAAAATATCAAGTGTTGGGAAAGAAAAAGGAAGTGGAATGGGTAACTCTTCTTGATTA AAAGTTATGTAATAACCAAATGCAATGTGAAATATTTTACTGGACTCTTTTGAAAAAC CATCTGTAAAAGACTGGGGGTGGGGGGGGGGCCAGCACGGTGGTGAGGCAGTTGAGAAAA GAGAAGGGTGTTGTAGTTTATAAAAGACTGTCTTAATTTGCATACTTAAGCATT<u>TAGG</u> <u>ATGTGGCAAAATGTTACAGAATCTAACTGGTGGACATGGCTGTTCATTGTACTGTTTTT</u> TCTATCTTCTATATGTTTAAAAGTATATAATAAAAATATTTAATTT

FIGURE 3B

#### EP 0 708 178 A1

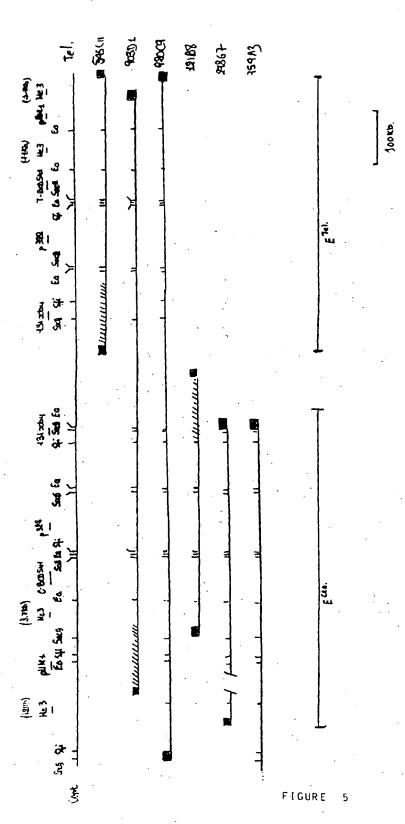
#### C212

#### C272

#### AFM157xd10

#### C161

### C171



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L-132										٠,					 											
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L-13							•								•				41	`.		_	_			
SMN pon	<b>10</b> -										5,"				 									3·	. •	

Telomeric element (ETel) containing the survival motor-neuron gene (SMN gene). Genetic map shows polymorphic markers C212, C272 and C171. Physical map shows location and direction of transcription of SMN gene; phage clones used for assembling physical map. Restriction map for EcoRI(E), XbaI(X), HindIII(H), BglII(B), SacII(S) are shown. Cent. and Tel. indicate centromere and telomere respectively. The position of genomic rearrangements found in SMA patients are also indicated.

Gane dosage analysis of the 5q13 region with the 132SE11 plasmid cone in SMA type I patient. Total human DNA from SMA family was digested with HindIII for Southern blotting. Filter was consecutively hybridized with 132SE11 (A) and JK53 probes (B). A significant decrease in 132SE11 band intensity, which indicated the deletion, compared with their parents. F/Father, M/Mother, A/affected

MAMSSGGSGGVPEQEDSVLFRRGTGQSDDSDIWDDTALIKAYDKAVASFKHA
LKNGDICETSGKPKTTPKRKPAKKNKSQKKNTAASLQQWKVGDKCSAIWSEDG
CIYPATIASIDFKRETCVVVYTGYGNREEQNLSDLLSPICEVANNIEQNAQEN
ENESQVSTDESENSRSPGNKSDNIKPKSAPWNSFLPPPPPMPGPRLGPGKPGL
KFNGPPPPPPPPPPHLLSCWLPPFPSGPPIIPPPPPICPDSLDDADALGSMLI
SWYMSGYHTGYYM



# EUROPEAN SEARCH REPORT

Application Number P 94 40 2353

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